## FIGURE 1

1 GAATTCAAGA CCAGCCTGGA CAACTTGGAA GAACCCGGTC TCTACAAAAA ATACAAAATT 61 AGCTGGGATT GGGTGCGGTG GCTCATGCCT ATAATCCCAG CACTTTGGGA GCCTGAGGTG 121 GGTGGATCAC CTGAAGTCAG GAGTTCAAGA CTAGCCTGGC CAACATGGTG AAACCCTATC 181 TCTACTGAAA ATACAAAAAG CTAGACGTGG TGGCACACAC CTGTAATCCC AGCTACTTAG 241 GAGGCTGAGG CAGGAGAATT GCTTGAAGCC TAGAGGTGAA GGTTGTAGTG AGCCGAGATT 301 GCATCATTGC ACAATGGAGG GGAGCCACCA GCCTGGGCAA CAAGAGGAAA TCTCCGTCTC 361 CAAAAAAAA AAAAAAAAA AAAGAATTAG GCTGGGTGGT GCCTGTAGTC CCAGCTACTT 421 GGGAGGCAGG GGGTCCACTT GATGTCGAGA CTGCAGTGAG CCATGATCCT GCCACTGCAC 481 TCCGGCCTGG GCAACAGAGT GAGACCCTGT CTAAAGAAAA AAAAAATAAA GCAACATATC 541 CTGAACAAG GATCCTCCAT AACGTTCCCA CCAGATTTCT AATCAGAAAC ATGGAGGCCA 601 GAAAGCAGTG GAGGAGGACG ACCCTCAGGC AGCCCGGGAG GATGTTGTCA CAGGCTGGGG 781 GGTCCCATCC AGGAAACCTC CGGCATGGCT GGGAAGTGGG GTACTTGGTG CCGGGTCTGT 841 ATGTGTGTGT GACTGGTGTG TGTGAGAGAG AATGTGTGCC CTAAGTGTCA GTGTGAGTCT 901 GTGTATGTGT GAATATTGTC TTTGTGTGGG TGATTTTCTG CGTGTGTAAT CGTGTCCCTG 961 CAAGTGTGAA CAAGTGGACA AGTGTCTGGG AGTGGACAAG AGATCTGTGC ACCATCAGGT 1021 GTGTGCATAG CGTCTGTGCA TGTCAAGAGT GCAAGGTGAA GTGAAGGGAC CAGGCCCATG 1081 ATGCCACTCA TCATCAGGAG CTCTAAGGCC CCAGGTAAGT GCCAGTGACA GATAAGGGTG 1141 CTGAAGGTCA CTCTGGAGTG GGCAGGTGGG GGTAGGGAAA GGGCAAGGCC ATGTTCTGGA 1201 GGAGGGGTTG TGACTACATT AGGGTGTATG AGCCTAGCTG GGAGGTGGAT GGCCGGGTCC 1261 ACTGAAACCC TGGTTATCCC AGAAGGCTTT GCAGGCTTCA GGAGCTTGGA GTGGGGAGAG 1321 GGGGTGACTT CTCCGACCAG GCCCCTCCAC CGGCCTACCC TGGGTAAGGG CCTGGAGCAG 1381 GAAGCAGGGG CAAGAACCTC TGGAGCAGCC CATACCCGCC CTGGCCTGAC TCTGCCACTG 1441 GCAGCACAGT CAACACAGCA GGTTCACTCA CAGCAGAGGG CAAAGGCCAT CATCAGCTCC 1501 CTTTATAAGG GAAGGGTCAC GCGCTCGGTG TGCTGAGAGT GTCCTGCCTG GTCCTCTGTG 1561 CCTGGTGGGG TGGGGGTGCC AGGTGTGTCC AGAGGAGCCC ATTTGGTAGT GAGGCAGGTA 1621 TGGGGCTAGA AGCACTGGTG CCCCTGGCCG TGATAGTGGC CATCTTCCTG CTCCTGGTGG 1681 ACCTGATGCA CCGGCGCCAA CGCTGGGCTG CACGCTACCC ACCAGGCCCC CTGCCACTGC
1741 CCGGGCTGGG CAACCTGCTG CATGTGGACT TCCAGAACAC ACCATACTGC TTCGACCAGG
1801 TGAGGGAGGA GGTCCTGGAG GGCGGCAGAG GTGCTGAGGC TCCCCTACCA GAAGCAAACA 1861 TGGATGGTGG GTGAAACCAC AGGCTGGACC AGAAGCCAGG CTGAGAAGGG GAAGCAGGTT 1921 TGGGGGACGT CCTGGAGAAG GGCATTTATA CATGGCATGA AGGACTGGAT TTTCCAAAGG 1981 CCAAGGAAGA GTAGGGCAAG GGCCTGGAGG TGGAGCTGGA CTTGGCAGTG GGCATGCAAG 2041 CCCATTGGGC AACATATGTT ATGGAGTACA AAGTCCCTTC TGCTGACACC AGAAGGAAAG 2101 GCCTTGGGAA TGGAAGATGA GTTAGTCCTG AGTGCCGTTT AAATCACGAA ATCGAGGATG 2161 AAGGGGGTGC AGTGACCCGG TTCAAACCTT TTGCACTGTG GGTCCTCGGG CCTCACTGCC 2221 TCACCGGCAT GGACCATCAT CTGGGAATGG GATGCTAACT GGGGCCTCTC GGCAATTTTG 2281 GTGACTCTTG CAAGGTCATA CCTGGGTGAC GCATCCAAAC TGAGTTCCTC CATCACAGAA 2341 GGTGTGACCC CCACCCCGC CCCACGATCA GGAGGCTGGG TCTCCTCCTT CCACCTGCTC 2401 ACTCCTGGTA GCCCCGGGGG TCGTCCAAGG TTCAAATAGG ACTAGGACCT GTAGTCTGGG 2461 GTGATCCTGG CTTGACAAGA GGCCCTGACC CTCCCTCTGC AGTTGCGGCG CCGCTTCGGG 2521 GACGTGTTCA GCCTGCAGCT GGCCTGGACG CCGGTGGTCG TGCTCAATGG GCTGGCGGCC 2581 GTGCGCGAGG CGCTGGTGAC CCACGGCGAG GACACCGCCG ACCGCCGCC TGTGCCCATC 2641 ACCCAGATCC TGGGTTTCGG GCCGCGTTCC CAAGGCAAGC AGCGGTGGGG ACAGAGACAG 2701 ATTTCCGTGG GACCCGGGTG GGTGATGACC GTAGTCCGAG CTGGGCAGAG AGGGCGCGGG 2761 GTCGTGGACA TGAAACAGGC CAGCGAGTGG GGACAGCGGG CCAAGAAACC ACCTGCACTA 2821 GGGAGGTGTG AGCATGGGGA CGAGGGCGGG GCTTGTGACG AGTGGGCGGG GCCACTGCCG 2881 AGACCTGGCA GGAGCCCAAT GGGTGAGCGT GGCGCATTTC CCAGCTGGAA TCCGGTGTCG 2941 AAGTGGGGGC GGGGACCGCA CCTGTGCTGT AAGCTCAGTG TGGGTGGCGC GGGGCCCGCG 3001 GGGTCTTCCC TGAGTGCAAA GGCGGTCAGG GTGGGCAGAG ACGAGGTGGG GCAAAGCCTG 3061 CCCCAGCCAA GGGAGCAAGG TGGATGCACA AAGAGTGGGC CCTGTGACCA GCTGGACAGA 3121 GCCAGGGACT GCGGGAGACC AGGGGGAGCA TAGGGTTGGA GTGGGTGGTG GATGGTGGGG 3181 CTAATGCCTT CATGGCCACG CGCACGTGCC CGTCCCACCC CCAGGGGTGT TCCTGGCGCG 3241 CTATGGGCCC GCGTGGCGCG AGCAGAGGCG CTTCTCCGTG TCCACCTTGC GCAACTTGGG 3301 CCTGGGCAAG AAGTCGCTGG AGCAGTGGGT GACCGAGGAG GCCGCCTGCC TTTGTGCCGC 3361 CTTCGCCAAC CACTCCGGTG GGTGATGGGC AGAAGGGCAC AAAGCGGGAA CTGGGAAGGC 3421 GGGGGACGGG GAAGGCGACC CCTTACCCGC ATCTCCCACC CCCAGGACGC CCCTTTCGCC 3481 CCAACGGTCT CTTGGACAAA GCCGTGAGCA ACGTGATCGC CTCCCTCACC TGCGGGCGCC 3541 GCTTCGAGTA CGACGACCCT CGCTTCCTCA GGCTGCTGGA CCTAGCTCAG GAGGGACTGA 3601 AGGAGGAGTC GGGCTTTCTG CGCGAGGTGC GGAGCGAGAG ACCGAGGAGT CTCTGCAGGG 3661 CGAGCTCCCG AGAGGTGCCG GGGCTGGACT GGGGCCTCGG AAGAGCAGGA TTTGCATAGA 3721 TGGGTTTGGG AAAGGACATT CCAGGAGACC CCACTGTAAG AAGGGCCTGG AGGAGGAGGG 3781 GACATCTCAG ACATGGTCGT GGGAGAGGTG TGCCCGGGTC AGGGGGCACC AGGAGAGGCC 3841 AAGGACTCTG TACCTCCTAT CCACGTCAGA GATTTCGATT TTAGGTTTCT CCTCTGGGCA 3901 AGGAGAGAG GTGGAGGCTG GCACTTGGGG AGGGACTTGG TGAGGTCAGT GGTAAGGACA

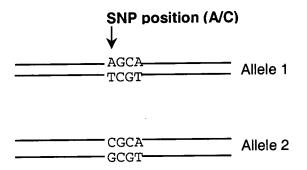
3961 GGCAGGCCCT GGGTCTACCT GGAGATGGCT GGGGCCTGAG ACTTGTCCAG GTGAACGCAG 4021 AGCACAGGAG GGATTGAGAC CCCGTTCTGT CTGGTGTAGG TGCTGAATGC TGTCCCCGTC 4081 CTCCTGCATA TCCCAGCGCT GGCTGGCAAG GTCCTACGCT TCCAAAAGGC TTTCCTGACC 4141 CAGCTGGATG AGCTGCTAAC TGAGCACAGG ATGACCTGGG ACCCAGCCCA GCCCCCCGA 4201 GACCTGACTG AGGCCTTCCT GGCAGAGATG GAGAAGGTGA GAGTGGCTGC CACGGTGGGG 4261 GGCAAGGGTG GTGGGTTGAG CGTCCCAGGA GGAATGAGGG GAGGCTGGGC AAAAGGTTGG 4321 ACCAGTGCAT CACCCGGCGA GCCGCATCTG GGCTGACAGG TGCAGAATTG GAGGTCATTT 4381 GGGGGCTACC CCGTTCTGTC CCGAGTATGC TCTCGGCCCT GCTCAGGCCA AGGGGAACCC 4441 TGAGAGCAGC TTCAATGATG AGAACCTGCG CATAGTGGTG GCTGACCTGT TCTCTGCCGG 4501 GATGGTGACC ACCTCGACCA CGCTGGCCTG GGGCCTCCTG CTCATGATCC TACATCCGGA 4561 TGTGCAGCGT GAGCCCATCT GGGAAACAGT GCAGGGGCCG AGGGAGGAAG GGTACAGGCG 4621 GGGGCCCATG AACTTTGCTG GGACACCCGG GGCTCCAAGC ACAGGCTTGA CCAGGATCCT 4681 GTAAGCCTGA CCTCCTCCAA CATAGGAGGC AAGAAGGAGT GTCAGGGCCG GACCCCCTGG 4741 GTGCTGACCC ATTGTGGGGA CGCATGTCTG TCCAGGCCGT GTCCAACAGG AGATCGACGA 4801 CGTGATAGGG CAGGTGCGGC GACCAGAGAT GGGTGACCAG GCTCACATGC CCTACACCAC 4861 TGCCGTGATT CATGAGGTGC AGCGCTTTGG GGACATCGTC CCCCTGGGTG TGACCCATAT 4921 GACATCCCGT GACATCGAAG TACAGGGCTT CCGCATCCCT AAGGTAGGCC TGGCGCCCTC 4981 CTCACCCAG CTCAGCACCA GCACCTGGTG ATAGCCCCAG CATGGCTACT GCCAGGTGGG 5041 CCCACTCTAG GAACCCTGGC CACCTAGTCC TCAATGCCAC CACACTGACT GTCCCCACTT 5101 GGGTGGGGG TCCAGAGTAT AGGCAGGGCT GGCCTGTCCA TCCAGAGCCC CCGTCTAGTG 5161 GGGAGACAAA CCAGGACCTG CCAGAATGTT GGAGGACCCA ACGCCTGCAG GGAGAGGGGG 5221 CAGTGTGGGT GCCTCTGAGA GGTGTGACTG CGCCCTGCTG TGGGGTCGGA GAGGGTACTG
5281 TGGAGCTTCT CGGGCGCAGG ACTAGTTGAC AGAGTCCAGC TGTGTGCCAG GCAGTGTGTG 5341 TCCCCGTGT GTTTGGTGGC AGGGGTCCCA GCATCCTAGA GTCCAGTCCC CACTCTCACC 5401 CTGCATCTCC TGCCCAGGGA ACGACACTCA TCACCAACCT GTCATCGGTG CTGAAGGATG 5461 AGGCCGTCTG GGAGAAGCCC TTCCGCTTCC ACCCCGAACA CTTCCTGGAT GCCCAGGGCC 5521 ACTTTGTGAA GCCGGAGGCC TTCCTGCCTT TCTCAGCAGG TGCCTGTGGG GAGCCCGGCT 5581 CCCTGTCCCC TTCCGTGGAG TCTTGCAGGG GTATCACCCA GGAGCCAGGC TCACTGACGC 5641 CCCTCCCCTC CCCACAGGCC GCCGTGCATG CCTCGGGGAG CCCCTGGCCC GCATGGAGCT 5701 CTTCCTCTC TTCACCTCCC TGCTGCAGCA CTTCAGCTTC TCGGTGCCCA CTGGACAGCC 5761 CCGGCCCAGC CACCATGGTG TCTTTGCTTT CCTGGTGAGC CCATCCCCT ATGAGCTTTG
5821 TGCTGTGCCC CGCTAGAATG GGGTACCTAG TCCCCAGCCT GCTCCCTAGC CAGAGGCTCT 5881 AATGTACAAT AAAGCAATGT GGTAGTTCCA ACTCGGGTCC CCTGCTCACG CCCTCGTTGG
5941 GATCATCCTC CTCAGGGCAA CCCCACCCCT GCCTCATTCC TGCTTACCCC ACCGCCTGGC 6001 CGCATTTGAG ACAGGGGTAC GTTGAGGCTG AGCAGATGTC AGTTACCCTT GCCCATAATC 6061 CCATGTCCCC CACTGACCCA ACTCTGACTG CCCAGATTGG TGACAAGGAC TACATTGTCC 6121 TGGCATGTGG GGAAGGGGCC AGAATGGGCT GACTAGAGGT GTCAGTCAGC CCTGGATGTG 6181 GTGGAGAGGG CAGGACTCAG CCTGGAGGCC CATATTTCAG GCCTAACTCA GCCCACCCCA 6241 CATCAGGGAC AGCAGTCCTG CCAGCACCAT CACAACAGTC ACCTCCCTTC ATATATGACA 6301 CCCCAAAACG GAAGACAAAT CATGGCGTCA GGGAGCTATA TGCCAGGGCT ACCTACCTCC 6361 CAGGGCTCAG TCGGCAGGTG CCAGAACGTT CCCTGGGAAG GCCCCATGGA AGCCCAGGAC 6421 TGAGCCACCA CCCTCAGCCT CGTCACCTCA CCACAGGACT GGCTACCTCT CTGGGCCCTC 6481 AGGGATGCTG CTGTACAGAC CCCTGACCAG TGACGAGTTC GCACTCAGGG CCAGGCTGGC 6541 GCTGGAGGA GACACTTGTT TGGCTCCAAC CCTAGGTACC ATCCTCCAG TAGGGATCAG
6601 GCAGGGCCCA CAGGCCTGCC CTAGGGACAG GAGTCAACCT TGGACCCATA AGGCACTGGG
6661 GCGGGCAGAG AAGGAGGAGG TGGCATGGGC AGCTGAGAGC CAGAGACCCT GACCCTAGTC
6721 CTTGCTCTGC CATTACCCCG TGTGACCCCG GGCCCACCCT TCCCCACCCC 6781 GGGCTTCTGT TTCCTTCTGC CAACGAGAAG GCTGCTTCAC CTGCCCCGAG TCCTGTCTTC 6841 CTGCTCTGCC TTCTGGGGCT GTGGCCCTTG CTGGCCTGGA GCCCCAACCA AGGGCAGGGA 6901 CTGCTGTCCT CCACGTCTGT CCTCACCGAC ATAATGGGCT GGGCTGGGCA CACAGGCAGT 6961 GCCCAAGAGT TTCTAATGAG CATATGATTA CCTGAGTCCT GGGCAGACCT TCTTAGGGAA 7021 CAGCCTGGGA CAGAGAACCA CAGACACTCT GAGGAGCCAC CCTGAGGCCT CTTTTGCCAG 7081 AGGACCCTAC AGCCTCCCTG GCAGCAGTTC CGCCAGCATT TCTGTAAATG CCCTCATGCC 7141 AGGGTGCGGC CCGGCTGTCA GCACGAGAGG GACGTTGGTC TGTCCCCTGG CACCGAGTCA 7201 GTCAGAAGGG TGGCCAGGGC CCCCTTGGGC CCCTCCAGAG ACAATCCACT GTGGTCACAC 7261 GGCTCGGTGG CAGGAAGTGC TGTTCCTGCA GCTGTGGGGA CAGGGAGTGT GGATGAAGCC 7321 AGGCTGGGTT TGTCTGAAGA CGGAGGCCCC GAAAGGTGGC AGCCTGGCCT ATAGCAGCAG 7381 CAACTCTTGG ATTTATTGGA AAGATTTTCT TCACGGTTCT GAGTCTTGGG GGTGTTAGAG 7441 GCTCAGAACC AGTCCAGCCA GAGCTCTGTC ATGGGCACGT AGACCCGGTC CCAGGGCCTT 7501 TGCTCTTTGC TGTCCTCAGA GGCCTCTGCA AAGTAGAAAC AGGCAGCCTT GTGAGTCCCC 7741 GAAGGTGGGA AAGGGCTGGG GTGTCTGTGA CCCTGGCAGT CACTGAGAAG CAGGGTGGAA 7801 GCAGCCCCT GCAGCACGCT GGGTCAGTGG TCTTACCAGA TGGATACGCA GCAACTTCCT 7861 TTTGAACCTT TTTATTTTCC TGGCAGGAAG AAGAGGGATC CAGCAGTGAG ATCAGGCAGG 7921 TTCTGTGTTG CACAGACAGG GAAACAGGCT CTGTCCACAC AAAGTCGGTG GGGCCAGGAT 7981 GAGGCCCAGT CTGTTCACAC ATGGCTGCTG CCTCTCAGCT CTGCACAGAC GTCCTCGCTC

8041	CCCTGGGATG	GCAGCTTGGC	CTGCTGGTCT	TGGGGTTGAG	CCAGCCTCCA	GCACTGCCTC
8101	CCTGCCCTGC	TGCCTCCCAC	TCTGCAGTGC	TCCATGGCTG	CTCAGTTGGA	CCCACGCTGG
8161	AGACGTTCAG	TCGAAGCCCC	GGGCTGTCCT	TACCTCCCAG	TCTGGGGTAC	CTGCCACCTC
8221	CTGCTCAGCA	GGAATGGGGC	TAGGTGCTTC	CTCCCCTGGG	GACTTCACCT	GCTCTCCCTC
8281	CTGGGATAAG	ACGGCAGCCT	CCTCCTTGGG	GGCAGCAGCA	TTCAGTCCTC	CAGGTCTCCT
8341	GGGGGTCGTG	ACCTGCAGGA	GGAATAAGAG	GGCAGACTGG	GCAGAAAGGC	CTTCAGAGCA
8401	CCTCATCCTC	CTGTTCTCAC	ACTGGGGTGT	CACAGTCCTG	GGAAGTTCTT	CCTTTTCAGT
8461	TGAGCTGTGG	TAACCTTGTG	AGTTTCCTGG	AGGGGGCCTG	CCACTACCCT	TGGGACTCCC
8521	TGCCGTGTGT	CTGGGTCTAA	CTGAGCTCTG	AAAGGAGAGA	GCCCCAGCCC	TGGGCCTTCC
8581	AGGGGAAGCC	TTACCTCAGA	GGTTGGCTTC	TTCCTACTCT	TGACTTTGCG	TCTCTGCAGA
8641	GGGAGGTGGG	AGGGGTGACA	CAACCCTGAC	ACCCACACTA	TGAGTGATGA	GTAGTCCTGC
8701	CCCGACTGGC	CCATCCTTTC	CAGGTGCAGT	CCCCCTTACT	GTGTCTGCCA	AGGGTGCCAG
	CACAGCCGCC	CCACTCCAGG	GGAAGAGGAG	TGCCAGCCCT	TACCACCTGA	GTGGGCACAG
8821	TGTAGCATTT	ATTCATTAGC	CCCCACACTG	GCCTGACCAT	CTCCCCTGTG	GGCTGCATGA
8881	CAAGGAGAGA	GAACAGGCTG	AGGTGAGAGC	TACTGTCAAC	ACCTAAACCT	AAAAAATCTA
	TAATTGGGCT	GGGCAGGGTG	GCTCACGCCT	GTAATCCCAG	CACTTTGGGA	GGCCGAGATG
	GGTGGATCAC	CTGAGGTCAG	ATGTTCGAGA	CCAGCCTGGC	CAACATGGTG	AAACCCCGTC
	TCTACTAAAA	ATACAAAAAA	TTAGCTGGGC	GTGGTGGTGG	GTGCCTGTAA	TCCCAGCTAC
	TCAGGAGGCT	GAGGCAGGAG	AATTGCTTGA	ACCTGGGAGG	CAGAGGCTGC	AGTGAGCCGA
	GATCGCATCA	TTGCACTCCA	GCCTGGTCAA	CAAGAGTGAA	ACTGTCTTAA	AAAAAAAATC
	TATAATTGAT	ATCTTTAGAA	AGATAAAACT	TTGCATTCAT	GAAATAAGAA	TAGGAGGGTC
	TAAAATAAAA	ATGTTCAAAC	ACCCACCACC	ACTAATTCTT	GACAAAAATA	TAGTCTGGGT
	GCCTTAGCTC	ATGCCTGTAA	TCCCAGCATT	TTGGGAGGCT	AAGGCAGGAG	GATTGTTTGA
9421	GCCTAGGAAT	TC				

## FIGURE 2

1	CAAMMCAACA	CCACCCTCCA	CAACMMCCAA	CAACCECCEC	TCTACAAAAA	**************************************
61				_		
					CACTTTGGGA	
121					CAACATGGTG	
181		_			CTGTAATCCC	
241	GAGGCTGAGG	CAGGAGAATT	GCTTGAAGCC	TAGAGGTGAA	GGTTGTAGTG	AGCCGAGATT
301	GCATCATTGC	ACAATGGAGG	GGAGCCACCA	GCCTGGGCAA	CAAGAGGAAA	TCTCCGTCTC
361	CAAAAAAAAA	AAAAAAAAA	AAAG <b>R</b> ATTAG	GCTGGGTGGT	GCCTGTAGTC	CCAGCTACTT
421	GGGAGGCAGG	GGGTCCACTT	GATGTCGAGA	CTGCAGTGAG	CCATGATCCT	GCCACTGCAC
481	TCCGGCCTGG	GCAACAGAGT	GAGACCCTGT	CTAAAGAAAA	AAAAAATAAA	GCAACATATC
541	CTGAACAAAG	GATCCTCCAT	AACGTTCCCA	CCAGATTTCT	AATCAGAAAC	ATGGAGGCCA
601	GAAAGCAGTG	${\tt GAGGAGGAC}\underline{{\tt R}}$	ACCCTCAGGC	AGCCCGGGAG	GATGTTGTCA	CAGGCTGGGG
661	CAAGGGCCTT	CCGGCTACCA	ACTGGGAGCT	CTGGGAACAG	CCCTGTTGCA	AACAAGAAGC
721	CATAGCCCGG	CCAGAGCCCA	GGAATGTGGG	CTGGGCTGGG	AGCAGCCTCT	GGACAGGAGT
781	GGTCCCATCC	AGGAAACCTC	CGGCATGGCT	GGGAAGTGGG	GTACTTGGTG	CCGGGTCTGT
841	ATGTGTGTGT	GACTGGTGTG	TGTGAGAGAG	$\mathtt{AATGTGTGC}\underline{\mathbf{Y}}$	CTAAGTGTCA	GTGTGAGTCT
901	GTGTATGTGT	GAATATTGTC	TTTGTGTGGG	TGATTTTCTG	CRTGTGTAAT	CGTGTCCCTG
961	CAAGTGTGAA	CAAGTGGACA	AGTGTCTGGG	AGTGGACAAG	AGATCTGTGC	ACCATCAGGT
1021	GTGTGCATAG	CGTCTGTGCA	TGTCAAGAGT	GCAAGGTGAA	GTGAAGGGAC	CAGGCCCATG
1081	ATGCCACTCA	TCATCAGGAG	CTCTAAGGCC	CCAGGTAAGT	GCCAGTGACA	GATAAGGGTG
1141	CTGAAGGTCA	CTCTGGAGTG	GGCAGGTGGG	GGTAGGGAAA	GGGCAAGGCC	ATGTTCTGGA
1201	GGAGGGGTTG	TGACTACATT	AGGGTGTATG	AGCCTAGCTG	GGAGGTGGAT	GGCC <b>R</b> GGTCC
1261	ACTGAAACCC	TGGTTATCCC	AGAAGGCTTT	GCAGGCTTCA	GGAGCTTGGA	GTGGGGAGAG
1321	GGGGTGACTT	CTCCGACCAG	GCCCCTCCAC	CGGCCTACCC	TGGGTAAGGG	CCTGGAGCAG
1381	GAAGCAGGGG	CAAGAACCTC	TGGAGCAGCC	CATACCCGCC	CTGGCCTGAC	TCTGCCACTG
1441	GCAGCACAGT	CAACACAGCA	GGTTCACTCA	CAGCAGAGGG	CAAAGGCCAT	CATCAGCTCC
1501	CTTTATAAGG	GAAGGGTCAC	GCGCTCGGTG	TGCTGAGAGT	GTCCTGCCTG	GTCCTCTGTG
1561	CCTGGTGGGG	TGGGGGTGCC	AGGTGTGTCC	AGAGGAGCCC	ATTTGGTAGT	GAGGCAGGTA
1621	${\tt TGGGGCTAGA}$	AGCACTGGTG	CCCCTGGCCG	TGATAGTGGC	CATCTTCCTG	CTCCTGGTGG

## FIGURE 3 One Base Sequencing (OBS) Outline



## Add Cy5-ddATP + dTTP,dCTP,dGTP + DNA polymerase

